



#3

SEQUENCE LISTING

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Mueller, Joerg P.
Van Dijl, Jan M.

<120> Twin-Arginine Translocation in Bacillus

<130> GC634-2

<140> US 09/954,737

<141> 2001-09-17

<150> US 60/233,610

<151> 2000-09-18

<160> 29

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<210> 1

<211> 89

<212> PRT

<213> Escherichia coli

<400> 1

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1 5 10 15
Val Leu Leu Phe Gly Thr Lys Lys Leu Gly Ser Ile Gly Ser Asp Leu
20 25 30
Gly Ala Ser Ile Lys Gly Phe Lys Lys Ala Met Ser Asp Asp Glu Pro
35 40 45
Lys Gln Asp Lys Thr Ser Gln Asp Ala Asp Phe Thr Ala Lys Thr Ile
50 55 60
Ala Asp Lys Gln Ala Asp Thr Asn Gln Glu Gln Ala Lys Thr Glu Asp
65 70 75 80
Ala Lys Arg His Asp Lys Glu Gln Val
85

<210> 2

<211> 67

<212> PRT

<213> Escherichia coli

<400> 2

Met Gly Glu Ile Ser Ile Thr Lys Leu Leu Val Val Ala Ala Leu Val
1 5 10 15
Val Leu Leu Phe Gly Thr Lys Lys Leu Arg Thr Leu Gly Gly Asp Leu
20 25 30
Gly Ala Ala Ile Lys Gly Phe Lys Lys Ala Met Asn Asp Asp Asp Ala
35 40 45
Ala Ala Lys Lys Gly Ala Asp Val Asp Leu Gln Ala Glu Lys Leu Ser
50 55 60
His Lys Glu
65

<210> 3

<211> 57
 <212> PRT
 <213> Bacillus subtilis

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 Met Pro Ile Gly Pro Gly Ser Leu Ala Val Ile Ala Ile Val Ala Leu
 1 5 10 15
 Ile Ile Phe Gly Pro Lys Lys Leu Pro Glu Leu Gly Lys Ala Ala Gly
 20 25 30
 Asp Thr Leu Arg Glu Phe Lys Asn Ala Thr Lys Gly Leu Thr Ser Asp
 35 40 45
 Glu Glu Glu Lys Lys Lys Glu Asp Gln
 50 55

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 <211> 70
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 <213> Bacillus subtilis

Q16
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 Met Phe Ser Asn Ile Gly Ile Pro Gly Leu Ile Leu Ile Phe Val Ile
 1 5 10 15
 Ala Ile Ile Ile Phe Gly Pro Ser Lys Leu Pro Glu Ile Gly Arg Ala
 20 25 30
 Ala Lys Arg Thr Leu Leu Glu Phe Lys Ser Ala Thr Lys Ser Leu Val
 35 40 45
 Ser Gly Asp Glu Lys Glu Glu Lys Ser Ala Glu Leu Thr Ala Val Lys
 50 55 60
 Gln Asp Lys Asn Ala Gly
 65 70

<210> 5
 <211> 62
 <212> PRT
 <213> Bacillus subtilis

<400> 5
 Met Glu Leu Ser Phe Thr Lys Ile Leu Val Ile Leu Phe Val Gly Phe
 1 5 10 15
 Leu Val Phe Gly Pro Asp Lys Leu Pro Ala Leu Gly Arg Ala Ala Gly
 20 25 30
 Lys Ala Leu Ser Glu Phe Lys Gln Ala Thr Ser Gly Leu Thr Gln Asp
 35 40 45
 Ile Arg Lys Asn Asp Ser Glu Asn Lys Glu Asp Lys Gln Met
 50 55 60

<210> 6
 <211> 171
 <212> PRT
 <213> Escherichia coli

<400> 6
 Met Phe Asp Ile Gly Phe Ser Glu Leu Leu Leu Val Phe Ile Ile Gly
 1 5 10 15
 Leu Val Val Leu Gly Pro Gln Arg Leu Pro Val Ala Val Lys Thr Val
 20 25 30
 Ala Gly Trp Ile Arg Ala Leu Arg Ser Leu Ala Thr Thr Val Gln Asn
 35 40 45
 Glu Leu Thr Gln Glu Leu Lys Leu Gln Glu Phe Gln Asp Ser Leu Lys
 50 55 60

Lys Val Glu Lys Ala Ser Leu Thr Asn Leu Thr Pro Glu Leu Lys Ala
 65 70 75 80
 Ser Met Asp Glu Leu Arg Gln Ala Ala Glu Ser Met Lys Arg Ser Tyr
 85 90 95
 Val Ala Asn Asp Pro Glu Lys Ala Ser Asp Glu Ala His Thr Ile His
 100 105 110
 Asn Pro Val Val Lys Asp Asn Glu Ala Ala His Glu Gly Val Thr Pro
 115 120 125
 Ala Ala Ala Gln Thr Gln Ala Ser Ser Pro Glu Gln Lys Pro Glu Thr
 130 135 140
 Thr Pro Glu Pro Val Val Lys Pro Ala Ala Asp Ala Glu Pro Lys Thr
 145 150 155 160
 Ala Ala Pro Ser Pro Ser Ser Ser Asp Lys Pro
 165 170

<210> 7
 <211> 258
 <212> PRT
 <213> Escherichia coli

916
 <400> 7
 Met Ser Val Glu Asp Thr Gln Pro Leu Ile Thr His Leu Ile Glu Leu
 1 5 10 15
 Arg Lys Arg Leu Leu Asn Cys Ile Ile Ala Val Ile Val Ile Phe Leu
 20 25 30
 Cys Leu Val Tyr Phe Ala Asn Asp Ile Tyr His Leu Val Ser Ala Pro
 35 40 45
 Leu Ile Lys Gln Leu Pro Gln Gly Ser Thr Met Ile Ala Thr Asp Val
 50 55 60
 Ala Ser Pro Phe Phe Thr Pro Ile Lys Leu Thr Phe Met Val Ser Leu
 65 70 75 80
 Ile Leu Ser Ala Pro Val Ile Leu Tyr Gln Val Trp Ala Phe Ile Ala
 85 90 95
 Pro Ala Leu Tyr Lys His Glu Arg Arg Leu Val Val Pro Leu Leu Val
 100 105 110
 Ser Ser Ser Leu Leu Phe Tyr Ile Gly Met Ala Phe Ala Tyr Phe Val
 115 120 125
 Val Phe Pro Leu Ala Phe Gly Phe Leu Ala Asn Thr Ala Pro Glu Gly
 130 135 140
 Val Gln Val Ser Thr Asp Ile Ala Ser Tyr Leu Ser Phe Val Met Ala
 145 150 155 160
 Leu Phe Met Ala Phe Gly Val Ser Phe Glu Val Pro Val Ala Ile Val
 165 170 175
 Leu Leu Cys Trp Met Gly Ile Thr Ser Pro Glu Asp Leu Arg Lys Lys
 180 185 190
 Arg Pro Tyr Val Leu Val Gly Ala Phe Val Val Gly Met Leu Leu Thr
 195 200 205
 Pro Pro Asp Val Phe Ser Gln Thr Leu Leu Ala Ile Pro Met Tyr Cys
 210 215 220
 Leu Phe Glu Ile Gly Val Phe Phe Ser Arg Phe Tyr Val Gly Lys Gly
 225 230 235 240
 Arg Asn Arg Glu Glu Glu Asn Asp Ala Glu Ala Glu Ser Glu Lys Thr
 245 250 255
 Glu Glu

<210> 8
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 <212> PRT
 <213> Bacillus subtilis

<400> 8

Met Thr Arg Met Lys Val Asn Gln Met Ser Leu Leu Glu His Ile Ala
 1 5 10 15
 Glu Leu Arg Lys Arg Leu Leu Ile Val Ala Leu Ala Phe Val Val Phe
 20 25 30
 Phe Ile Ala Gly Phe Phe Leu Ala Lys Pro Ile Ile Val Tyr Leu Gln
 35 40 45
 Glu Thr Asp Glu Ala Lys Gln Leu Thr Leu Asn Ala Phe Asn Leu Thr
 50 55 60
 Asp Pro Leu Tyr Val Phe Met Gln Phe Ala Phe Ile Ile Gly Ile Val
 65 70 75 80
 Leu Thr Ser Pro Val Ile Leu Tyr Gln Leu Trp Ala Phe Val Ser Pro
 85 90 95
 Gly Leu Tyr Glu Lys Glu Arg Lys Val Thr Leu Ser Tyr Ile Pro Val
 100 105 110
 Ser Ile Leu Leu Phe Leu Ala Gly Leu Ser Phe Ser Tyr Tyr Ile Leu
 115 120 125
 Phe Pro Phe Val Val Asp Phe Met Lys Arg Ile Ser Gln Asp Leu Asn
 130 135 140
 Val Asn Gln Val Ile Gly Ile Asn Glu Tyr Phe His Phe Leu Leu Gln
 145 150 155 160
 Leu Thr Ile Pro Phe Gly Leu Leu Phe Gln Met Pro Val Ile Leu Met
 165 170 175
 Phe Leu Thr Arg Leu Gly Ile Val Thr Pro Met Phe Leu Ala Lys Ile
 180 185 190
 Arg Lys Tyr Ala Tyr Phe Thr Leu Leu Val Ile Ala Ala Leu Ile Thr
 195 200 205
 Pro Pro Glu Leu Leu Ser His Met Met Val Thr Val Pro Leu Leu Ile
 210 215 220
 Leu Tyr Glu Ile Ser Ile Leu Ile Ser Lys Ala Ala Tyr Arg Lys Ala
 225 230 235 240
 Gln Lys Ser Ser Ala Ala Asp Arg Asp Val Ser Ser Gly Gln
 245 250

<210> 9

<211> 245

<212> PRT

<213> Bacillus subtilis

<400> 9

Met Asp Lys Lys Glu Thr His Leu Ile Gly His Leu Glu Glu Leu Arg
 1 5 10 15
 Arg Arg Ile Ile Val Thr Leu Ala Ala Phe Phe Leu Phe Leu Ile Thr
 20 25 30
 Ala Phe Leu Phe Val Gln Asp Ile Tyr Asp Trp Leu Ile Arg Asp Leu
 35 40 45
 Asp Gly Lys Leu Ala Val Leu Gly Pro Ser Glu Ile Leu Trp Val Tyr
 50 55 60
 Met Met Leu Ser Gly Ile Cys Ala Ile Ala Ala Ser Ile Pro Val Ala
 65 70 75 80
 Ala Tyr Gln Leu Trp Arg Phe Val Ala Pro Ala Leu Thr Lys Thr Glu
 85 90 95
 Arg Lys Val Thr Ile Met Tyr Ile Met Tyr Ile Pro Gly Leu Phe Ala
 100 105 110
 Leu Phe Leu Ala Gly Ile Ser Phe Gly Tyr Phe Val Leu Phe Pro Ile
 115 120 125
 Val Leu Ser Phe Leu Thr His Leu Ser Ser Gly His Phe Glu Thr Met
 130 135 140
 Phe Thr Ala Asp Arg Tyr Phe Arg Phe Met Val Asn Leu Ser Leu Pro

145 150 155 160
 Phe Gly Phe Leu Phe Glu Met Pro Leu Val Val Met Phe Leu Thr Arg
 165 170 175
 Leu Gly Ile Leu Asn Pro Tyr Arg Leu Ala Lys Ala Arg Lys Leu Ser
 180 185 190
 Tyr Phe Leu Leu Ile Val Val Ser Ile Leu Ile Thr Pro Pro Asp Phe
 195 200 205
 Ile Ser Asp Phe Leu Val Met Ile Pro Leu Leu Val Leu Phe Glu Val
 210 215 220
 Ser Val Thr Leu Ser Ala Phe Val Tyr Lys Lys Arg Met Arg Glu Glu
 225 230 235 240
 Thr Ala Ala Ala Ala
 245

<210> 10
 <211> 63
 <212> PRT
 <213> Bacillus alcalophilus

<400> 10
 Met Gly Gly Leu Ser Val Gly Ser Val Val Leu Ile Ala Leu Val Ala
 1 5 10 15
 Leu Leu Ile Phe Gly Pro Lys Lys Leu Pro Glu Leu Gly Lys Ala Ala
 20 25 30
 Gly Ser Thr Leu Arg Glu Phe Lys Asn Ala Thr Lys Gly Leu Ala Asp
 35 40 45
 Asp Asp Asp Asp Thr Lys Ser Thr Asn Val Gln Lys Glu Lys Ala
 50 55 60

<210> 11
 <211> 272
 <212> PRT
 <213> Bacillus alcalophilus

<400> 11
 Met Thr Met Met Thr Pro Asn Gln Gln Thr Ser Lys Lys Lys Lys Arg
 1 5 10 15
 Lys Gly Arg Lys Gly Arg Val Pro Met Gln Asp Met Ser Ile Met Asp
 20 25 30
 His Ala Glu Leu Arg Arg Arg Ile Phe Val Val Leu Ala Phe Phe
 35 40 45
 Ile Val Ala Leu Ile Gly Gly Phe Phe Leu Ala Val Pro Val Ile Thr
 50 55 60
 Phe Leu Gln Asn Ser Pro Gln Ala Ala Asp Met Pro Phe Asn Ala Phe
 65 70 75 80
 Arg Leu Thr Asp Pro Leu Arg Val Tyr Met Asn Phe Ala Val Ile Thr
 85 90 95
 Ala Leu Val Leu Ile Ile Pro Val Ile Leu Tyr Gln Leu Trp Ala Phe
 100 105 110
 Val Ser Pro Gly Leu Lys Glu Asn Glu Gln Lys Ala Thr Leu Ala Tyr
 115 120 125
 Ile Pro Ile Ala Phe Leu Leu Phe Leu Ala Gly Ile Ala Phe Ser Tyr
 130 135 140
 Phe Ile Leu Leu Pro Phe Val Ile Ser Phe Met Gly Gln Met Ala Asp
 145 150 155 160
 Arg Leu Glu Ile Asn Glu Met Tyr Gly Ile Asn Glu Tyr Phe Ser Phe
 165 170 175
 Leu Phe Gln Leu Thr Ile Pro Phe Gly Leu Leu Phe Gln Leu Pro Val
 180 185 190
 Val Val Met Phe Leu Thr Arg Leu Gly Val Val Thr Pro Thr Phe Leu

195 200 205
 Arg Lys Ile Arg Lys Tyr Ala Tyr Phe Ala Leu Leu Val Ile Ala Gly
 210 215 220
 Ile Ile Thr Pro Pro Glu Leu Thr Ser His Leu Phe Val Thr Val Pro
 225 230 235 240
 Met Leu Ile Leu Tyr Glu Ile Ser Ile Thr Ile Ser Ala Ile Thr Tyr
 245 250 255
 Arg Lys Tyr His Gly Thr Thr Asp His Asn Gly Gln Glu Ser Ala Lys
 260 265 270

<210> 12
 <211> 35
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 12
 cccaagctta tgaaaggag ggcttttttg aatgg

35

<210> 13
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 13
 gcggatccaa agctgagcac gatcgg

26

<210> 14
 <211> 39
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 14
 cccaagctta aaaagaaaga agatcagtaa gttaggatg

39

<210> 15
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer

<400> 15
 gcggatccaa gtcctgagaa atccg

25

<210> 16
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>

<223> primer

<400> 16

ggaattcgtg ggacggctac c

21

<210> 17

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 17

cgggatccat catgggaagc g

21

<210> 18

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 18

ggggtaccgg aaaacgcttg atcagg

26

<210> 19

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 19

cgggatcctt tgggcgatag cc

22

<210> 20

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 20

gaggatccat gaggagagag gggatcttga atggcatacg ac

42

<210> 21

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 21

cgatcctgca ggacctcatc ggattgc

27

<210> 22
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<220>
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<400> 22
gtaggatccg cgcctaactt ctcaagc

27

<210> 23
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<220>
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<400> 23
atagaattca aaaaggaaga gtatg

25

<210> 24
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<400> 24
ctggggatcc aaaaacagga aggc

24

<210> 25
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<400> 25
gagaaggatcg acgcagcatt tacttcaaag gcccc

35

<210> 26
<211> 26
<212> DNA
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<220>
<223> primer

<400> 26
accgggtcga ccgctgtttt acaacg

26

<210> 27
<211> 23
<212> DNA
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<220>
<223> primer

<400> 27
gggaattcat ggcctgcccg gtt

23

<210> 28
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 28
caaggatccc gaattaagga gtgg

24

<210> 29
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
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<400> 29
ggtctgcagc tgcactaagc ggccgcc

27

916
ccc